

Figure 1

SEQ ID NO:25	***** ** *** ** ** ** **	MVASS-QKILVTSAGFIGTHTVVQLLNNGNFVSIIDNFDNSVMEAVREVVGSNLSQ	60
SEQ ID NO:26		MSS---QTVLVTGGAGYIGSHTVLQLLGGFKAVVDNLDNSSETAIHRVKELAGK-FAG	
SEQ ID NO:14		MVSSS-QHILVTGGAGFIGTHTVVQLLKAGFSVSIIDNFDNSVMEAVDRVQVVGPLLSQ	
SEQ ID NO:16		--AR-GSVLVTGGAGFIGTHTVLQLLEKGYAVTAVDNFHNVSPEALDRVRHIVGPALSA	
SEQ ID NO:18		T-----	
SEQ ID NO:20		MVSALLRTILVTGGAGYIGSHTVLQLLQLGFRVVLDNLDNASELAILRVRELAGEH-NAN	
SEQ ID NO:22		MRD---KTVLVTGGAGYIGSHTVLQLLGGFRAVVLDNLDNLENSESSEVAIHRVRELAGE-FGN	1
			60
SEQ ID NO:25	* * * * *	NLEFTLGLRNKDDLEKLFSSKSKFDVAIHFAGLKAVGESVENPRRYFDNNLVGTINLYEV	
SEQ ID NO:26		NLSFHKLDLRDRDALEKIFSSSTKSDSVIHFAGLKAVGESVQKPLLYDNNLIGTIVLFEV	
SEQ ID NO:14		NLQFTQGLNRDDLEKLFSKTTFDVAIHFAGLKAVAESVAKPRRYFDNVLGTINLYEF	
SEQ ID NO:16		RLQFIFGDLTIKDDLEKVFAAKKYDAVIHFAGLKAVAESVAHPMYNNRNNIVGTINLYDV	
SEQ ID NO:18		-----RIDLRDKGALEMVFASTRFEAVIHFAGLKAVGESVQKPLLYDNNVIGTINLLEV	
SEQ ID NO:20		NLDFRKVDLRDKQALDQIFSSQRFEAVIHFAGLKAVGESVQKPLLYDNNLIGTITLLQV	
SEQ ID NO:22		NLSFHKVDLRDRAALDQIFSSSTQFDVAIHFAGLKAVGESVQKPLLYNNNLTGTITLLEV	61
			120
SEQ ID NO:25	* * * * *	MAKHNCCKMVFSSSATVYGQPEKIPCVEDFKLQAMNPYGRGTKLFLEEIARDIQKAEPEWR	
SEQ ID NO:26		MAAHGCKKLVFSSSATVYGLPKVPCTEEFPLSAANPYGRTKLIIEEICRDIYRAEQEWK	
SEQ ID NO:14		MAKYNCKKMVFSSSATVYGQPEKIPCEEDFKLQAMNPYGRTKLFLEEIARDIQKAEPEWK	
SEQ ID NO:16		MKKHGCNKLVFSSSATVYGQPEKVPCEFDSPKALNPYGRTKLYLEEMLRDYQHANPEWR	
SEQ ID NO:18		MSVHGCCKLVFSSSAAVYGPSKNSPCTENFPLTPNNPYGKTKLVVEDICRDIYRSDPEWK	
SEQ ID NO:20		MAAHGCTKLVFSSSATVYGWPKVPCTEESPLCAMNPYGRTKLVIEDMCRDLHASDPNWK	
SEQ ID NO:22		MAAHGCKKLVFSSSATVYGWPKVPCTEEFPLSAMNPYGRTKLIIEEICRDVHCAEPDCK	121
			180

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Figure 1

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***** *** ** ***** * ***** * ***** * * * * * * * * *
SEQ ID NO:25 IVLLRYFNPVGAHESGKLGEDPRGIPNNLMPYIQQVAVAGRLPELNVYGHDPYTRDGSAIR
SEQ ID NO:26 IILLRYFNPVGAHPSGYIGEDPRGIPNNLMPFVQQVAVAGRRPALTVFGNDYTTSDGTGVR
SEQ ID NO:14 IILLRYFNPVGAHESGKLGEDPKGIPNNLMPYIQQVAVAGRLTELNVYGHDPYTRDGSAIR
SEQ ID NO:16 TILLRYFNPIGAHESGDI GEDPKGVPNNLLPYIQQVAVARRPELNVYGHDPYTRDGTAVR
SEQ ID NO:18 IILLRYFNPVGAHPSGYLGEDPRGIPNNLMPYVQQVAVAGRRPALTVLGNDYATRDGTGVR
SEQ ID NO:20 IILLRYFNPVGAHPSGYIGEDPCGIPNNLMPFVQQVAVAGRRPALTVYGTDYNTKDGTVR
SEQ ID NO:22 IILLRYFNPVGAHPSGYIGEDPRGIPNNLMPFVQQVAVAGRRPALTVFGNDYNTSDGTGVR
181
***** ***** **
SEQ ID NO:25 DYIHVMDLADGHIAALRKLFTSEN-IGCTAYNLGTRGSSVLEMVAAFEKASGKKIALKL
SEQ ID NO:26 DYIHVVDLADGHIAALRKL--NDPKIGCEVYNLGTGKGTSVLEMVKAFAQSAGKKIPLVM
SEQ ID NO:14 DYIHVMDLADGHIAALRKLFTTEN-IGCTAYNLGTRGTSVLEMVTAFAKASGKKIPVKL
SEQ ID NO:16 DYIHVVDLADGHIAALEKLFATPD-IGCVAYNLGTRGTTVLEMVSAFEKAYGKKIPVKM
SEQ ID NO:18 DYIHVVDLADGHIAALQKLFENSS-IGCEAYNLGTRGTSVLEIVKAFEKASGKKIPLIF
SEQ ID NO:20 DYIHVVDLADGHIAALRKLIEDSDRIGCEVYNLGTGKGTSVLEMVAAFEKASGKKIPLVF
SEQ ID NO:22 DYIHVVDLADGHIAALLKL--DEPNIGCEVYNLGTGKGTSVLEMVRAFEMASGKKIPLVM
241
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SEQ ID NO:25 CPRRPGDATEVYASTAKAEKELGWKAKYGV EEMCRDQWNWAKNNPWGYSG-----KP---
SEQ ID NO:26 AGRRPGDAEVVYASTNKAERELNWKAKYGIDEMCRDQWNWASKNPNPYGGSEDSSN-----
SEQ ID NO:14 CPRRPGDATEVYASTERAELGWKANYGV EEMCRDQWNWAKNNPWGYSAG-----KP---
SEQ ID NO:16 CPRRPGDSEQVYASTAKAEELGWRAKYGIEEMCRDQWNWAKNNPNPYGCGNAAENKD---
SEQ ID NO:18 GERRPGDAEILFSETTKAERELNWKAKYGIEEMCRDQWNWASKNPNPYGSPDSIKQNGHQ
SEQ ID NO:20 AGRRPGDAEIVYAQTAKAEKELWKAKYGV EEMCRDLWNWASKNPNPYGYG-SPDSSN-----
SEQ ID NO:22 AGRRPGDAEIVYASTKKAERELKWKAKYGIDEMCRDQWNWASKNPNPYGYGD-QGSTD-----
301

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Figure 1

SEQ ID NO:25	-----	
SEQ ID NO:26	-----	
SEQ ID NO:14	-----	
SEQ ID NO:16	-----	
SEQ ID NO:18	TNGSADSSKQNGHRTNGSTDSPKRNHAYGSADSPKRNHCVFGSSDLKPNGNGLR	
SEQ ID NO:20	-----	
SEQ ID NO:22	-----	
		361
		418